

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
27 June 2002 (27.06.2002)

PCT

(10) International Publication Number
WO 02/50304 A2

(51) International Patent Classification⁷: C12Q 1/04, A61P 35/00

(21) International Application Number: PCT/US01/45108

(22) International Filing Date: 3 December 2001 (03.12.2001)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:
60/250,131 1 December 2000 (01.12.2000) US
60/327,016 5 October 2001 (05.10.2001) US

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(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

— without international search report and to be republished upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

WO 02/50304 A2

(54) Title: ONCOLYTIC VIRUS

(57) Abstract: Methods of reducing the viability of a tumor cell, infecting a neoplasm in a mammal with a virus, or treating a neoplasm in a mammal, utilizing certain non-naturally occurring viruses are disclosed. Viral reassortants, for example reovirus reassortants, and techniques for identifying PKR-sensitive viruses are also disclosed.

ONCOLYTIC VIRUS

SUMMARY OF THE INVENTION

This invention provides methods of reducing the viability of a tumor cell, infecting a neoplasm in a mammal with a virus, or treating a neoplasm in a mammal, comprising administering a non-naturally occurring virus wherein the virus is: a) a reovirus whose mu-2 protein has amino acid residues A, R, M, F, L, M, I, Q, I and S at positions 93, 150, 300, 302, 347, 372, 434, 458, 652 and 726, respectively; or b) a reassortant of two or more parent strains of a viral species selected from the family Reoviridae, or progeny thereof; or c) a virus other than a reovirus wherein the virus other than a reovirus is: i) capable of expressing a reovirus mu-2 protein having amino acid residues A, R, M, F, L, M, I, Q, I and S at positions 93, 150, 300, 302, 347, 372, 434, 458, 652 and 726, respectively, and ii) is a DNA virus, a positive-sense RNA virus, or a negative-sense RNA virus selected from the group consisting of Orthomyxoviridae, Rhabdoviridae and Paramyxoviridae. This invention further provides the use of such non-naturally occurring virus in the manufacture of a medicament for reducing the viability of a tumor cell, infecting a neoplasm in a mammal, or treating a neoplasm in a mammal.

This invention provides a method of identifying a PKR sensitive virus comprising: a) dividing a sample of a virus to be tested into a first portion and second portion; b) contacting PKR +/+ cells with the first portion and contacting PKR -/- cells with the second portion, under conditions permitting growth of the virus in PKR -/- cells; c) determining the rate of growth of the virus in the PKR +/+ cells and in the PKR -/- cells; and d) comparing the growth rates from step c), wherein a higher rate of growth in the PKR -/- cells than in the PKR +/+ cells identifies the virus as PKR sensitive. Such PKR sensitive viruses identified in accordance with this invention are useful for reducing the viability of a tumor cell, infecting a neoplasm in a mammal, or treating a neoplasm in a mammal.

DESCRIPTION OF THE FIGURES

Figure 1: Virus yield of reovirus strains T1L and T3D in PKR -/- vs. PKR +/- murine embryo fibroblasts.

Figure 2: Immuno-blot of PKR in MEF Infected with Reo T1L and T3D.

Figure 3: Lungs of mice with ct26 tumors after treatment with reovirus strains. T1L, T3D, EB96, EB108 and EB146 relative to untreated control lung. The lungs from 2 mice are shown for each treatment.

Figure 4: The weight of BALB-C mouse lungs relative to the presence of CT26 tumors and reovirus treatment.

Figure 5: Histological sections stained with hematoxylin and eosin showing lung lobes of mice with ct26 tumors after treatment with reovirus strains. T1L, T3D, EB96, EB108 and EB146 relative to untreated control lung.

DETAILED DESCRIPTION OF THE INVENTION

Throughout this application amino acids are generally identified using the standard one-letter abbreviation, but can also be identified by name or standard three-letter abbreviations.

T3D, T1L, T3A and T2J are standard abbreviations for reovirus strains T3 Dearing, T1 Lang, T3 Abney, and T2 Jones, respectively. The above-listed names of strains and their respective abbreviations are used interchangeably.

As used herein "phenotype" refers to the sequence of the expressed proteins of a virus. In the case of reoviruses the expressed proteins are the gene products of the L1, L2, L3, M1, M2, M3, S1, S2, S3 and S4 genes. Thus, if the amino acid sequences of the products of these genes are the same in two different reoviral strains they are said to have the same phenotype.

As used herein "genotype" refers to the nucleotide sequence of the coding region of a virus. Thus, for example, if the nucleotide sequences of the L1, L2, L3, M1, M2, M3, S1, S2, S3

and S4 genes of two reoviruses are the same in two different reoviral strains they are said to have the same genotype.

The term "PFU" stands for plaque forming units and is a quantitative measure of live virus particles.

Examples of the anti-neoplastic and anti-tumor methods and use of this invention as described above, include those utilizing a reovirus whose mu-2 protein has amino acid residues A, R, M, F, L, M, I, Q, I and S at positions 93, 150, 300, 302, 347, 372, 434, 458, 652 and 726, respectively. In a more specific embodiment the reoviral mu-2 protein has the amino acid sequence of the mu-2 protein of reovirus strain T3 Dearing, for example when the mu-2 protein is expressed by a gene having the nucleic acid sequence of the M1 gene of reovirus strain T3 Dearing. In a more specific embodiment the reovirus has the same genotype as a reovirus strain selected from the group consisting of eb86, eb129, eb88, eb13, and eb145. In a more specific embodiment the reovirus has a M1 gene whose sequence is the same as the M1 gene of reovirus strain T3 Dearing and an L3 gene whose sequence is the same as the L3 gene of reovirus strain T1 Lang, for example the virus can have the same genotype as a reovirus strain selected from the group consisting of eb28, eb31, eb97, eb123 and g16. In a still more specific embodiment the reovirus has a M1 gene whose sequence is the same as the M1 gene of reovirus strain T3 Dearing and an L3 gene, L1 gene, and S2 gene whose sequences are the same as the corresponding genes of reovirus strain T1 Lang, for example reoviruses having the same genotype as a reovirus strain selected from eb96, eb146 and eb108. In an even more specific embodiment the reovirus has a M1 gene whose sequence is the same as the M1 gene of reovirus strain T3 Dearing and an L3 gene, L1 gene, S2 gene and S4 gene whose sequences are the same as the corresponding genes of reovirus strain T1 Lang, for example reoviruses having the same genotype as reovirus strain eb96.

Other examples of the anti-neoplastic and anti-tumor methods and use of this invention as described above, include those utilizing a virus that is a reassortant of two or more parent strains of a viral species selected from the family Reoviridae, or progeny thereof. For example, reassortants can be made of two, three or four of the reovirus strains T3 Dearing, T1 Lang, T3 Abney, and T2 Jones. In a more specific embodiment the reassortants are generated from parent strains T3 Dearing and T1 Lang. Examples of such strains include eb118, eb73.1, h17, h15, eb39, and h60 as well as the other strains shown in Tables 1 and 2.

Other examples of the anti-neoplastic and anti-tumor methods and use of this invention as described above, include those utilizing a virus other than a reovirus that is: i) capable of expressing a reovirus mu-2 protein having amino acid residues A, R, M, F, L, M, I, Q, I and S at positions 93, 150, 300, 302, 347, 372, 434, 458, 652 and 726, respectively, and ii) is a DNA virus, a positive-sense RNA virus, or a negative-sense RNA virus selected from the group consisting of the families Orthomyxoviridae, Rhabdoviridae and Paramyxoviridae. Examples of suitable DNA viruses include a Herpesvirus, Adenovirus, Parvovirus, Papovavirus, Iridovirus, Hepadenavirus, Poxvirus, mumps virus, human parainfluenza virus, measles virus or rubella virus. Examples of suitable a positive-sense RNA viruses include a Togavirus, Flavivirus, Picornavirus, or Coronavirus. Examples of suitable negative-sense RNA virus selected from the group consisting of Orthomyxoviridae, Rhabdoviridae and Paramyxoviridae include an influenza virus or a vesicular stomatitis virus.

In accordance with the method of identifying a PKR sensitive virus of this invention as described above, any PKR +/+ and -/- cells can be used, and the rate of growth of the virus is determined by any standard technique for monitoring viral growth including those that measure the number of virus particles directly or the quantity of viral proteins. In a specific embodiment the PKR cells are mouse embryo fibroblasts. In another specific embodiment the rate of growth of the virus is determined by a technique selected from the group consisting of plaque titer assay, antibody assay, and Western blot. Each of these techniques is exemplified below. Preferably the growth rate of the virus in PKR -/- cells is at least ten times higher than the growth rate in PKR +/+ cells.

In all of the anti-neoplastic and anti-tumor methods and use of this invention as described above, the virus can be a replication competent virus and/or a clonal virus. The virus can be administered by any conventional route, including but not limited to intranasally, intratracheally, intravenously, intraperitoneally or intratumorally. In accordance with the method or use of reducing the viability of a tumor cell described above, the virus can be administered to the tumor cell either *in vivo* or *ex vivo*. When the virus is administered to a mammal, the mammal can be either a human or a non-human mammal such as a mouse, sheep, cow, pig, dog or rabbit. While the optimal dose is expected to differ somewhat from patient to patient and can readily be determined by a skilled clinician, a dosage of from 3 x 10⁷ to 3 x 10⁹ PFU/kg is typical.

The viruses utilized in accordance with this invention can be produced by any conventional means, including reassortment among two or more parent virus strains or the use of standard recombinant genetic techniques. Once produced, such viruses can be reproduced by culturing in cells to produce progeny. The construction of reassortants of viruses is well known and is described, for example in Brown, et al., "The L2 Gene of Reovirus Serotype 3 Controls the Capacity to Interfere, Accumulate Deletions and Establish Persistent Infection" in Double-Stranded RNA Viruses, Compans, et al. eds. Elsevier (1983). For example, reassortants can be made of two, three or four of the reovirus strains T3 Dearing, T1 Lang, T3 Abney, and T2 Jones. Reassortants of T3 Dearing and T1 Lang are described in Example 2. Preferably the virus is replication competent and/or a clonal virus.

This invention will be better understood by reference to the following examples, which illustrate but are not intended to limit the invention described herein.

EXPERIMENTS

Experiment 1: Growth of Reovirus Strains T1L and T3D in PKR Knock-Out and Wild Type Fibroblast Cells

Viral Growth

The effect of PKR on reovirus infection was examined using PKR knock-out (PKR -/-) murine embryo fibroblasts (MEF). Both reovirus T1L and T3D grow to several fold higher titre in PKR -/- relative to PKR +/+ MEF, as measured by plaque assay. (Figure 1) This was associated with a higher percentage of antigen positive cells detected by fluorescent antibody staining described below. Consistent with this, infection of PKR -/- MEF resulted in several fold greater amounts of viral protein as assayed by western blot described below. Although both T1L and T3D grew to higher titres in cells lacking the PKR gene T1L virus grew to higher titres than T3D in either PKR -/- or PKR +/+ cells. (Figure 1)

Indirect Immunostaining

Cells were grown on glass coverslips in 35 mm diameter dishes and were infected with reovirus T1L or T3D at a multiplicity of infection (moi) of 10. After 48 hours incubation the cells were rinsed in PBS and fixed in prechilled acetone for 5 min. After rinsing in PBS (3x5min), 100 μ l of an appropriate dilution of type-specific rabbit antivirus antisera was applied and incubated at room temperature for 30 min. The coverslips were then rinsed in PBS (3x5min) and treated with the appropriate dilution of Cy3-conjugated donkey anti-rabbit antibody (Jackson ImmunoResearch Laboratories, Inc.) as the secondary antibody. After another 30 min incubation period at room temperature the coverslips were rinsed in PBS (3x5min) and mounted on glass slides in Gel/Mount (Biomedica Corp). All antibody dilutions were done in PBS/3 % BSA.

The samples were examined with a Zeiss microscope equipped with epifluorescence and a 40X 1.40 NA PlanApo objective. The images were collected using Image One Metamorph software and a Hamamatsu chilled charge-coupled digital camera (model C5985). Configuration of the digital images was done using Corel Presentations software.

Immunoblotting

Monolayer cultures of MEF were infected at a moi=10 with T1L or T3D virus as described above. At various times the culture medium was removed and the cells were rinsed with PBS before solubilizing in 1 ml of sample buffer (62.5mM Tris-HCl pH6.8, 10% glycerol, 2 % SDS, 0.05% bromophenol blue and 5 % 2-mercaptoethanol)(Laemmli). Aliquots of 25 ul volume were subjected to SDS PAGE and transblotted onto an Immobilon P membrane (Millipore) at 25V overnight at 4°C. The dried membrane was blocked with 5% (w/v) skim milk powder in PBS for 1hr at RT. This was followed by the addition of type specific rabbit anti-reovirus immune serum as the primary antibody in fresh blocking solution and incubation for 2hr at 4°C. The membrane was then washed three times in PBS and once in TBS (100 mM Tris Hcl pH 7.4, 0.9 % NaCl)to remove phosphate and incubated in 5% milk in TBS containing 1 ug/ml protein A conjugated to alkaline phoshatase obtained from Sigma Chemicals (Oakville, Ont) Finally the membrane was washed 4x in TBS before reaction with chromogenic substrate, nitro blue tetrazolium (NBT) (33 ug/ml) plus 5-bromo-4-chloro-3-indolyl phoshate (BCIP) (3.3 ul/ml), in alkaline phosphatase buffer (100mM NaCl, 5mM MgCl₂ and 100mM Tris-HCl pH9.5). The reaction was stopped with PBS containing 20mM EDTA.

Experiment 2: Reassortants Between Reovirus Strains T1L and T3D

Production of Genetic reassortants between Reovirus Serotype 1 Lang strain and Serotype 3 Dearing strain.

Mouse L929 cells were coinfecte with Reovirus Serotype 1 Lang strain (T1L) and Serotype 3 Dearing strain (T3D) at a multiplicity of infection of 5 each. Virus was harvested 24 hr post infection by 3 cycles of freezing and thawing before progeny viruses were isolated by 2 cycles of plaque isolation in L929 monolayers. Since each of the corresponding genome segments of T1L and T3D is distinguishable by electrophoretic mobility the genetic composition of each virus was determined by polyacrylamide gel electrophoresis of the segmented double stranded RNA (dsRNA) genome where the mobility of each segment is compared to the parental strains. Gels prepared as described by Laemmli contained 10% polyacrylamide and 0.27% methylene bis-acrylamide. Double-stranded RNA was obtained

from L929 cells infected for 3 days and solubilised in buffer containing sodium dodecyl sulphate and was detected in gels stained with ethidium bromide as described previously (Zou S. and E.G. Brown. (1992) Identification of Sequence elements containing signals for replication and encapsidation of the reovirus M1 genome segment. *Virology* 186:377-88.. The use of this panel of reassortants was first described by E.G. Brown, M. L. Nibert and B.N. Fields (1983) The L2 gene of reovirus serotype 3 controls the capacity to interfere, accumulate deletions and establish persistent infection. in Double-Stranded RNA Viruses. R.W. Compans and D.H.L. Bishop eds. Elsevier Science Publishing Co.

Growth of Reovirus

T1L, T3D and virus stocks from the reassortment procedure described above were prepared in L929 cells grown in Earl's Minimal Essential Medium (MEM) supplemented with 5 % fetal bovine serum and penicillin to 100 units/ml and streptomycin to 100 ug/ml until cytopathic effect was complete. Cells and culture supernatant were subjected to 3 cycles of freezing and thawing before titration by plaque assay.

Yields in Mouse Embryo Fibroblasts

Wild type PKR^{+/+} cells were obtained from Balb-C mice and PKR^{-/-} cells were obtained from PKR knockout mice. Cell cultures were produced using 15-17 days embryos that had been disaggregated by mincing and trypsin treatment. Cell monolayers were grown in 35 mm plastic dishes in MEM supplemented with 10% FBS and P/S at 37 C in a 5% CO₂ atmosphere. Cells were infected with titrated T1L, T3D or reassortant reovirus at a multiplicity of infection (moi) of 10 by adsorption of stock virus for 0.5 hr with agitation at 15 minute intervals. Unadsorbed virus was removed by 3 washes with 2 ml of warm PBS each before the addition of 3 ml of MEM supplemented with 5 % fetal bovine serum and penicillin to 100 units/ml and streptomycin to 100 ug/ml. The yield of T1L and T3D was assayed at time points over a 4 day period and is shown in Figure 1. Comparison of yields of virus from MEF cells infected with reassortant reovirus was done after 3 days incubation by plaque assay of duplicate cultures. The results are shown below in Table 1 (PKR^{-/-}) and Table 2 (PKR^{+/+}).

Plaque assay of reovirus in L929 Cells

Monolayer cultures of L929 cells were decanted of medium and infected in duplicate with 0.1 ml volumes of serially diluted virus in PBS. Virus was adsorbed for 0.5 hr before the application of 3 ml of MEM supplemented with 1 % agar, 5 % FBS and P/S. Cultures were incubated at 37 C and supplementary overlays of 2 ml aliquots of the same medium was added 3 and 6 days post infection. After 8 days of infection the monolayers were stained for 24 hr with 2 ml of the same overlay solution supplemented with neutral red (0.01 % weight/volume) to observe plaques.

Discussion

The genetic basis for the increased ability of T1L to grow in each cell type was determined using T1L x T3D reassortants. The difference in yield in wild type MEF (PKR +/+) segregated primarily with the M1 gene whereas the difference in yield in PKR -/- MEF was associated with the L1, L3, M3 and S2 genes and did not involve the M1 gene. The comparison of the genetic basis for replication in PKR +/+ relative to PKR -/- MEF cells indicates that the ability of the PKR gene to inhibit reovirus infection is dependent on the properties of the M1 gene. Furthermore the extent of replication and thus exploitation of PKR -/- cells is dependent on the nature of the L1, L3, M3 and S2 genes. Thus the reassortant viruses with the greatest differential ability to replicate in PKR -/- relative to PKR +/+ cells possess the T3D M1 gene and the viruses with the greatest ability to replicate in PKR -/- cells (characteristic of many tumor cells) possess the L1, L3, M3 and S2 genes of T1L. Such viruses are restricted in replication of PKR +/+ cells but replicate to a greater extent than either T1L or T3D in PKR -/- cells and are embodied in the properties of the reassortants eb96 and eb108. Statistical analyses of the experimental results are shown in Tables 1, 2 and 3.

The amino acid sequences of the T1L and T3D mu2 proteins are shown in Table 4. Each protein is 736 amino acids long and they differ at 10 aa positions. The observed difference in sensitivity to PKR seen as an ability to replicate in PKR+/+ relative to PKR-/- MEF cells is attributed to the difference in amino acid sequence between these proteins and thus M1 proteins of reoviruses with these amino acid changes or other substitutions at these positions are addressed herein. The mu2 protein is encoded by the M1 gene. The nucleotide

sequences of the T1L and T3D M1 gene are shown in Table 5. Each genome segment is 2304 nucleotides long and they differ at 51 nucleotide positions.

TABLE 1: PKR -/-

VIRUS	TITRE	L1	L2	L3	M1	M2	M3	S1	S2	S3	S4	RANK
eb146	7.00E+08	L	L	D	L	L	L	L	L	D	D	1
eb28	5.80E+08	D	D	D	D	D	D	D	D	D	D	2
eb108	4.70E+08	L	D	D	L	L	L	D	D	D	D	3
eb118	4.50E+08	D	D	L	D	D	D	D	L	L	L	4
T1L	4.30E+08	L	L	L	L	L	L	L	L	L	L	5
eb73.1	3.50E+08	L	D	L	D	D	D	D	D	D	D	6
eb31	3.20E+08	L	L	D	L	L	L	D	D	D	D	7
h17	3.00E+08	D	D	L	D	D	D	D	D	D	D	8
H15	2.80E+08	L	D	D	D	D	D	D	D	D	D	9
eb39	2.60E+08	L	D	D	D	D	D	D	D	D	D	10
eb96	1.80E+08	L	D	D	L	L	L	D	D	D	D	11
eb97	1.40E+08	D	D	D	D	D	D	D	D	D	D	12
h60	1.30E+08	D	D	L	D	D	D	D	D	D	D	13
T3D	1.20E+08	D	D	D	D	D	D	D	D	D	D	14
eb123	9.50E+07	D	D	L	D	D	D	D	D	D	D	15
g16	9.30E+07	L	L	D	L	D	D	D	L	L	L	16
eb86	8.50E+07	L	D	D	D	D	D	D	D	D	D	17
eb129	6.30E+07	D	D	D	D	D	D	D	L	L	D	18
eb88	6.00E+07	D	D	D	D	D	D	D	D	D	D	19
eb13	5.30E+07	D	D	D	D	D	D	D	D	D	L	20
eb145	1.30E+07	D	D	D	D	D	D	L	D	D	D	21
t-test	0.045	0.19	0.019	0.024	0.25	0.75	0.57	0.087	0.62	0.76		
M-W test	0.085	0.19	0.007	0.109	0.28	1	0.26	0.047	0.61	0.97		

TABLE 2: PKR +/+ (wild type)

VIRUS	TITRE	L1	L2	L3	M1	M2	M3	S1	S2	S3	S4	RANK
h60	3.96E+08	D	D	L	L	D	D	D	D	D	L	1
eb39	2.35E+08	L	D	D	L	D	D	D	D	D	D	2
H15	1.78E+08	L	D	D	L	D	D	D	D	D	L	3
eb118	1.76E+08	D	D	L	L	D	D	D	D	L	L	4
eb146	1.68E+08	L	L	L	D	L	L	L	L	L	D	5
T1L	1.50E+08	L	L	L	L	L	L	L	L	L	L	6
h17	1.46E+08	D	D	L	L	D	D	L	D	D	L	7
eb28	1.30E+08	D	D	L	D	D	D	D	L	D	D	8
eb73.1	1.23E+08	L	D	L	L	D	D	D	D	D	D	9
eb31	5.20E+07	L	L	L	D	L	L	L	D	D	L	10
eb123	4.88E+07	D	D	L	D	D	D	D	D	L	D	11
g16	4.03E+07	L	L	L	D	L	L	L	D	L	L	12
eb129	3.78E+07	D	D	D	D	D	L	D	L	L	D	13
eb97	2.35E+07	D	D	L	D	D	D	D	D	D	L	14
eb96	2.20E+07	L	D	L	D	L	L	L	L	D	L	15
eb108	1.33E+07	L	D	L	D	L	L	L	L	D	D	16
T3D	1.20E+07	D	D	D	D	D	D	D	D	D	D	17
eb13	7.50E+06	D	D	D	D	D	D	D	D	D	L	18
eb86	6.40E+06	L	D	D	D	D	L	D	D	D	L	19
eb88	6.00E+06	D	D	D	D	L	D	D	D	D	D	20
eb145	2.25E+06	D	D	D	D	D	L	L	D	D	D	21
t-test												
M-W test												

In Tables 1 and 2, parental origin of genome segments is indicated by L (T1L) or D (T3D). Statistical significance was determined using the t-test and the Mann-Whitney (MW) test.

TABLE 3: SUSCEPTIBILITY TO PKR SEGREGATES WITH THE M1 GENE

Gene	Single gene regression (R ² %)		Stepwise regression (R ² %)	
	PKR+/+	PKR-/-	PKR+/+	PKR-/-
L1	0	19 (P=.048)	0	L3 + L1 48 (P=.003)
L3	23.8 (P=.025)	36 (P=.004)	M1+L3 67.0 (P<.001)	36 (P=.004)
M1	51.6 (P<.001)	0	51.6 (P=<.001)	L3 + L1+ M1 56 (P=.0025)
S2	0	16 (P=.073)	0	L3+L1+M3+S2 63.4 (P<.001)

TABLE 4: Alignment of T1L (GenBank Accession No. CAA42570.1) and T3D (GenBank Accession No. AAA47256.1) mu2 proteins. These amino acid sequences were deduced from cDNA. Each protein is 736 nucleotides long and differs at 10 aa positions.

T1L	1	MAYIAVPAVVDSRSSEAIGLLESFGVDAGADANDVSYQDHDYVLDQLQYMLDGYEAGDVI	60
Consensus		MAYIAVPAVVDSRSSEAIGLLESFGVDAGADANDVSYQDHDYVLDQLQYMLDGYEAGDVI	
T3D	1	MAYIAVPAVVDSRSSEAIGLLESFGVDAGADANDVSYQDHDYVLDQLQYMLDGYEAGDVI	60
T1L	61	DALVHKNWLHHSVYCLLPPKSQQLLEYWKSNPSPVI	120
Consensus		DALVHKNWLHHSVYCLLPPKSQQLLEYWKSNPSPS	
T3D	61	I	
T1L	61	PDVNDRRLRKRLMLKKDLRKDDEYNQ	
Consensus			
T3D	61	DALVHKNWLHHSVYCLLPPKSQQLLEYWKSNPSPAI	
T1L	121	DALVHKNWLHHSVYCLLPPKSQQLLEYWKSNPSPAI	120
Consensus		PDVNDRRLRKRLMLKKDLRKDDEYNQ	
T3D	121	PDVNDRRLRKRLMLKKDLRKDDEYNQ	
T1L	121	LARAFKISDVYAPLISSTTSPMTMIQNLNQGEIVYTTTDRVIGARILLYAPRKYYASTLS	180
Consensus		LARAFKISDVYAPLISSTTSPMTMIQNLN	
T3D	121	GEIVYTTTDRVIGARILLYAPRKYYASTLS	
T1L	181	GEIVYTTTDRVIGARILLYAPRKYYASTLS	
Consensus		LARAFKISDVYAPLISSTTSPMTMIQNLNRGEIVYTTTDRVIGARILLYAPRKYYASTLS	
T3D	181	180	
T1L	181	FTMTKCIIPFGKEVGRVPHSRFNVGTFPSIATPKCFVMSGVDIESIPNEFIKLFYQRVKS	240
Consensus		FTMTKCIIPFGKEVGRVPHSRFNVGTFPSIATPKCFVMSGVDIESIPNEFIKLFYQRVKS	
T3D	181	FTMTKCIIPFGKEVGRVPHSRFNVGTFPSIATPKCFVMSGVDIESIPNEFIKLFYQRVKS	240
T1L	241	FTMTKCIIPFGKEVGRVPHSRFNVGTFPSIATPKCFVMSGVDIESIPNEFIKLFYQRVKS	
Consensus		VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	300
T3D	241	VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	
T1L	241	VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	
T1L	301	VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	
Consensus		VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	
T3D	301	VHANILNDISPQIVSDMINRKRLRVHTPSDRRAAQLMHPYHVKGASHVDVYKVDDVV	
T1L	301	LLEVVVDVADGLRNVSRKLTMHTVPVCILEMLGIEIADYCIRQEDGMFTDWFLLLTMLSDG	360
Consensus		L	
T3D	301	EVVDVADGLRNVSRKLTMHTVPVCILEMLGIEIADYCIRQEDGM	
T1L	301	TDWFLLLTMLSDG	
T1L	361	TDWFLLLTMLSDG	
Consensus		L	
T3D	361	FEVVVDVADGLRNVSRKLTMHTVPVCILEMLGIEIADYCIRQEDGMLTDWFLLLTMLSDG	360
T1L	361	FEVVVDVADGLRNVSRKLTMHTVPVCILEMLGIEIADYCIRQEDGMLTDWFLLLTMLSDG	
T1L	361	LTDERRTHCQYLINPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	420
Consensus		LTDERRTHCQYL	
T3D	361	NPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	
T1L	361	NPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	
T1L	421	LTDERRTHCQYLMNPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	420
Consensus		LTDERRTHCQYLMNPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	
T3D	421	LTDERRTHCQYLMNPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	
T1L	421	LTDERRTHCQYLMNPSSVPPDVILNISITGFINRHTIDVMPDIYDFVKPIGAVLPKGSFKS	
T1L	481	TIMRVLDISIILGVQIMPRAHVVDSDDEVGEQMEPTFEAVMEIYKGIAVGVDSDLDDLIKWV	480
Consensus		TIMRVLDISIILG	
T3D	481	QIMPRAHVVDSDDEVGEQMEPTFE AVMEIYKGIAVGVDSDLDDLIKWV	
T1L	481	QIMPRAHVVDSDDEVGEQMEPTFE AVMEIYKGIAVGVDSDLDDLIKWV	
T1L	481	TIMRVLDISIILGIQIMPRAHVVDSDDEVGEQMEPTFEQAVMEIYKGIAVGVDSDLDDLIKWV	480
Consensus		TIMRVLDISIILGIQIMPRAHVVDSDDEVGEQMEPTFEQAVMEIYKGIAVGVDSDLDDLIKWV	
T3D	481	TIMRVLDISIILGIQIMPRAHVVDSDDEVGEQMEPTFEQAVMEIYKGIAVGVDSDLDDLIKWV	480
T1L	481	TIMRVLDISIILGIQIMPRAHVVDSDDEVGEQMEPTFEQAVMEIYKGIAVGVDSDLDDLIKWV	
T1L	541	LNSDLIPHDDRLGQLFQAFLPLAKDLLAPMARKFYDNMSSEGRLLTFAHADSELLNANYF	540
Consensus		LNSDLIPHDDRLGQLFQAFLPLAKDLLAPMARKFYDNMSSEGRLLTFAHADSELLNANYF	
T3D	541	LNSDLIPHDDRLGQLFQAFLPLAKDLLAPMARKFYDNMSSEGRLLTFAHADSELLNANYF	540
T1L	541	LNSDLIPHDDRLGQLFQAFLPLAKDLLAPMARKFYDNMSSEGRLLTFAHADSELLNANYF	
T1L	541	GHLLRLKIPYITEVNLMIRKNREGGELFQLVLSYLYKMYATSAQPKWFGSLLRLICPWL	600

Consensus GHLLRLKIPYITEVNL MIRKNREGGELFQLVLSYLYKMYATSAQPKWFGSLLRLLICPWL
T3D 541 GHLLRLKIPYITEVNL MIRKNREGGELFQLVLSYLYKMYATSAQPKWFGSLLRLLICPWL 600

T1L 601 HMEKLIGEADPASTSAEIGWHIPREQLMQDGWC GCEDGFIPYVSIRAPRLV MEELMEKNW 660
consensus HMEKLIGEADPASTSAEIGWHIPREQLMQDGWC GCEDGFIPYVSIRAPRLV EELMEKNW
T3D 601 HMEKLIGEADPASTSAEIGWHIPREQLMQDGWC GCEDGFIPYVSIRAPRLV EELMEKNW 660

T1L 661 GQYHAQVIVTDQLVVGEPRRVSAKAVIKGNHLPVKLVSRFACFTLTAKYEMRLSCGHSTG 720
Consensus GQYHAQVIVTDQLVVGEPRRVSAKAVIKGNHLPVKLVSRFACFTLTAKYEMRLSCGHSTG
T3D 661 GQYHAQVIVTDQLVVGEPRRVSAKAVIKGNHLPVKLVSRFACFTLTAKYEMRLSCGHSTG 720

T1L 721 RGAAYNARLAFRSDL A 736
Consensus RGAAY ARALAFRSDL A
T3D 721 RGAAY SARLAFRSDL A 736

TABLE 5: Alignment of the nucleotide sequences of the T1L (GenBank Accession No. X59945.1) and T3D (GenBank Accession No M27261.1) M1 cDNA encoding mu-2 protein. The complete coding sequences are shown. Since reoviruses are double-stranded RNA viruses, the reoviral genome would contain "u" in place to "t". Each genome segment shown below is 2304 nucleotides long that differ at 51 nucleotide positions.

T1L 1	gctattcgccgtcatggcttacatcgcagttcctgcgggtggattcacgttcaagtga 60
T3D 1	gctattcgccgtcatggcttacatcgcagttcctgcgggtggattcacgttcgagtga 60
T1L 61	ggctattggactgctagaatcgttggagtagacgctggggctgatgcgaatgacgttc 120
T3D 61	ggctattggactgctagaatcgttggagtagacgctggggctgacgcgaatgacgttc 120
T1L 121	atatcaagatcatgactatgtgttggatcagttacagtatgttagatggatatgaggc 180
T3D 121	atatcaagatcatgactatgtgttggatcagttacagtacatgttagatggatatgaggc 180
T1L 181	tggcgacgttatcgatgcactcgccacaagaattggttacatcactccgtctattgctt 240
T3D 181	tggtgacgttatcgatgcactcgccacaagaattggttacatcactctgtctattgctt 240
T1L 241	gttgcacccaaaagtcaactactagagtattggaaaagtaatccttcagtgataccgga 300
T3D 241	gttgcacccaaaagtcaactattagagtattggaaaagtaatccttcagcgataccgga 300
T1L 301	caacgttgcgtcggttcgtaaacgactaatgctaaagaaaagatctcagaaaagatga 360
T3D 301	caacgttgcgtcggttcgtaaacgactaatgctaaagaaaagatctcaggaaaagatga 360

T1L 361	tgaatacaatcaactagcgcgtgcttcaagatatcgatgtctacgcacctctcatctc	420
T3D 361	tgaatacaatcagctagcgcgtgcttcaagatatcgatgtctacgcacctctcatctc	420
T1L 421	atccacgacgtcaccgatgacaatgatccagaacttgaatcaaggcgagatcgtgtacac	480
T3D 421	atccacgacgtcaccgatgacaatgatacagaacttgaatcgaggcgagatcgtgtacac	480
T1L 481	cacgacggacaggtaattggggtagaatcttgttatgtctctagaaaagtactatgc	540
T3D 481	cacgacggacaggtaataggggtagaatcttgttatgtctctagaaaagtactatgc	540
T1L 541	gtcaactctatcatttactatgactaagtgcattccgttggcaaagaggtgggtcg	600
T3D 541	gtcaactctgtcatttactatgactaagtgcattccgttggtaaagaggtgggtcg	600
T1L 601	tgttcctcactctagatttaatgtggcacattccatcaattgtcacccgaaatgttt	660
T3D 601	tgttcctcactctcgatttaatgtggcacattccgtcaattgtcacccgaaatgttt	660
T1L 661	tgtcatgagtgggttgatattgagtcacatccaaatgaattcatcaagttgtttacca	720
T3D 661	tgtcatgagtgggttgatattgagtcacatccaaatgaatttatcaagttgtttacca	720
T1L 721	gcgctcaagagtgttacgccaatataactaaatgacatataccctcagatcgtctctga	780
T3D 721	gcgctcaagagtgttacgctaacataactaaatgacatataccctcagatcgtctctga	780
T1L 781	catgataaacagaaagcgccccatccatcgatcgtcgagccgcgcagtt	840
T3D 781	catgataaacagaaagcgctcgccgttccatcgatcgtcgagccgcgcagtt	840
T1L 841	gatgcatttgccttaccatgttaaacgaggagcgacgtcacgtcgacgtttacaagggtgga	900
T3D 841	gatgcatttgccttaccatgttaaacgaggagcgacgtcacgtcgacgtttacaagggtgga	900

T1L 901 tggtagacgttttagaggttgtggatgtggccatgggtgcgcaacgtatctag 960
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 901 tggtagacatgttgcaggtagtgtggatgtggccatgggtgcgcaacgtatctag 960

T1L 961 gaaactaactatgcataccgtccgtatgtattcttgaatgtgggtattgagattgc 1020
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 961 gaaactaactatgcataccgtccgtatgtattcttgaatgtgggtattgagattgc 1020

T1L 1021 ggactattgcattcgtcaagaggatggaatgtcacagattggccactttaccat 1080
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 1021 ggactattgcattcgtcaagaggatgctcacagattggccactttaccat 1080

T1L 1081 gctatctgtggcttaactgtatagaaggacgcattgtcaataacttgatccgtcaag 1140
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 1081 gctatctgtggcttgcactgtatagaaggacgcattgtcaataacttgatccgtcaag 1140

T1L 1141 tgtgcctcctgtatgtgatacttaacatctcaattactggatttataaataggcatacaat 1200
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 1141 tgtgcctcctgtatgtgatacttaacatctcaattactggatttataaatagacatacaat 1200

T1L 1201 cgatgtcatgcctgatatatgtacttcgttaaacccattggcgctgtgcctaagg 1260
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
T3D 1201 cgatgtcatgcctgacatatatgtacttcgttaaacccattggcgctgtgcctaagg 1260

T1L 1261	atcatttaaatcaacaatttatgagagttcttgcattcaatatcaatattaggagtcacat 1320
T3D 1261	atcatttaaatcaacaatttatgagagttcttgcattcaatatcaatattaggaaatccaaat 1320
T1L 1321	catgccgcgcgcgcatgttagttgactcgatgagggtggcgagcaaatggagcctacgtt 1380
T3D 1321	catgccgcgcgcatgttagttgactcagatgagggtggcgagcaaatggagcctacgtt 1380
T1L 1381	tgagcatgcggtatggagatatacaaaggattgctggcgttgcactcgctggatgatct 1440
T3D 1381	tgagcaggcggttatggagatatacaaaggattgctggcgttgcactcgctggatgatct 1440
T1L 1441	catcaagtgggtgctgaactcgatctcattccgcatgtgcacaggcttggccaattatt 1500
T3D 1441	catcaagtgggtgttgcactcgatctcattccgcatgtgcacaggcttggcaattatt 1500
T1L 1501	tcaagcgtttgcctctcgcaaaggacttgttagctccaatggccagaaagtttatga 1560
T3D 1501	tcaagcgttttgcctctcgcaaaggactttagctccaatggccagaaagtttatga 1560
T1L 1561	taactcaatgagtgggttagattgcgtacattcgctcatgccacagtgcgtttttatga 1620
T3D 1561	taactcaatgagtgggttagattgcgtacattcgctcatgccacagtgcgtttttatga 1620
T1L 1621	cgcaaattttggtcattttgcgactaaaaataccatattacagaggtaatct 1680
T3D 1621	cgcaaattttggtcattttgcgactaaaaataccatattacagaggtaatct 1680
T1L 1681	gatgattcgcaagaatcgtgggtggagagctattcagctgtgttatcgatctata 1740
T3D 1681	gatgattcgcaagaatcgtgggtggagagctattcagctgtgttatcttatctata 1740
T1L 1741	taaaaatgttatgtactagcgccgcagccataatggttggatcattattgcgtttatgttaat 1800
T3D 1741	taaaaatgttatgtactagcgccgcagccataatggttggatcattattgcgtttatgttaat 1800

T1L 1801 atgtccctggttacatatggagaaattaataggagaaggcagacccggcatctacgtcggc 1860
||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
T3D 1801 atgtccctggttacatatggagaaattaataggagaaggcagacccggcatctacgtcggc 1860

T1L 1861 tgaaaattggatggcatatccctcgtaacagctgatgcaagatggatggatgtga 1920
||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||
T3D 1861 tgaaaattggatggcatatccctcgtaacagctgatgcaagatggatggatgtga 1920

T1L 1921 agatggattcattccctatgttagcatacgtgcgccaagactggttatggaggagttgat 1980
||| | ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
T3D 1921 agacggattcattccctatgttagcatacgtgcgccaagactggttatagaggagttgat 1980

T1L 1981 ggagaagaactggggccaatatcatgcccaagttattgtcactgatcagttgtcgtagg 2040
||||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
T3D 1981 ggagaagaactggggccaatatcatgcccaagttattgtcactgatcagttgtcgtagg 2040

T1L 2041 cgaaccgcggagggtatctgccaaggctgtgatcaaggtaatcacttaccagttaaagt 2100
||||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
T3D 2041 cgaaccgcggagggtatctgctaaggctgtgatcaaggtaaccacttaccagttaaagt 2100

T1L 2101 agtttcacgatttgcattcacattgacggcgaagtatgagatgaggctctcgtaggg 2160
||||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
T3D 2101 agtttcacgatttgcattcacattgacggcgaagtatgagatgaggcttcgtcgtaggg 2160

T1L 2161 ccatagcactggacggggggctgcataacaatgcgagactagcttccgatctgacttggc 2220
||||||| ||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
T3D 2161 ccatagcactggacgtggagctgcatacagtgcgagactagcttccgatctgacttggc 2220

T1L 2221 gtgatccgtgacatgcgttagtgtgacacctgcccctaggtaatggggtaggggcggg 2280
||||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
T3D 2221 gtgatccgtgacatgcgttagtgtgacacctgctctaggtaatggggtaggggcggg 2280

T1L 2281 ctaagactacgtacgcgttcatc 2304
||||||| |||||
T3D 2281 ctaagactacgtacgcgttcatc 2304

Experiment 3: Assessment of lethal infection in PKR -/- vs. PKR +/+ Mice

Adult Balb-C PKR +/+ or PKR -/- mice were infected with various dosages of infectious reovirus T1L or T3D via the intraperitoneal (IP) or intranasal (IN) route. IP injections involved the administration of 0.1 ml of stock virus or virus diluted in PBS. IN infection involved the application of 0.05 ml volumes of stock virus or virus diluted in PBS onto the nose-pad of mice anaesthetized with halothane (administered at 3% in oxygen). The survival of adult mice was monitored over a 30 day period. Adult PKR +/+ and PKR -/- mice resisted infection with 5e6 infectious T3D virus whereas T1L virus killed PKR -/- mice but not PKR +/+ mice at this dose. This demonstrates an enhanced ability of T1L to infect the tissues of PKR -/- mice. Table 5.

Two day old suckling Balb-C PKR +/+ or PKR -/- mice were infected with various dosages of infectious reovirus T1L or T3D via the IN route. IN infection involved the application of 0.01 ml volumes of stock virus or virus diluted in PBS onto the nose-pad of mice anaesthetized with halothane (administered at 3% in oxygen). The survival of suckling mice was monitored over an 18 day period. Suckling PKR +/+ or PKR -/- mice were both susceptible to similar dosages of T1L whereas T3D virus killed PKR -/- mice much more effectively than PKR +/+ mice, killing them at doses more than 100 fold less than those required to kill wild type suckling mice. This demonstrates an enhanced ability of T3D to infect the tissues of PKR -/- tissues of suckling mice and indicates a difference in the properties of the T1L and the T3D strains with respect to differential replication in PKR +/+ versus PKR -/- mice although both viruses were more restricted in replication of PKR +/+ mice of different ages (adult versus suckling). Table 5.

TABLE 5

ADULT MICE	T1L virus (S/So)		T3D virus (S/So)	
	PKR+/+	PKR-/-	PKR+/+	PKR-/-
5 E6 IP	ND	100 % (3/3)	ND	100 % (3/3)
5 E6 IN	100 % (3/3)	0 % (0/3)	100 % (3/3)	100 % (3/3)
5 E5 IN	ND	100 % (3/3)	ND	ND
SUCKLING MICE				
	PKR+/+	PKR-/-	PKR+/+	PKR-/-
3 E6 IN	33 % (2/6)	66 % (2/3)	84 % (5/6)	0 % (0/2)
3 E4 IN	100 % (7/7)	ND	100 % (7/7)	0 % (0/4)
3 E3 IN	ND	ND	ND	100 % (3/3)

Experiment 4: Reovirus T3D is a stronger inducer of PKR MEF than T1L

Infection of PKR^{+/+} MEF results in a greater expression of the phosphorylated form of PKR (Fig. 2). PKR^{+/+} MEF were infected at a moi of 10 and incubated over a 48 hr period for immunoblot analysis using rabbit anti-PKR serum that reacts with the first 100 amino acids of PKR. Proteins were separated on a 10% polyacrylamide gel and transferred to IMMOBILON membrane (Millipore Inc.) before incubation with 1/100 diluted primary antibody in the presence of casein. After repeated washing the blot was incubated with goat anti-rabbit antibody conjugated with alkaline phosphatase (1/30,000 dilution) (Sigma Inc) for 1 hour before repeated washing and reaction with Attophos substrate for phosphorescent detection as shown in Figure 2. Activation of PKR results in an electrophoretic form of slightly slower mobility indicated as PKR-P. Infection with T3D results in a greater production of this form than with infection with T1L. This demonstrates that PKR expression is enhanced in T3D infected cells and indicates that this may be responsible for the greater sensitivity of this virus to the PKR gene.

Experiment 5: Proof of principle for Improved Oncolysis of reovirus T1L x T3D Reassortants: Demonstration that reovirus reassortants with the M1 gene of T3D and the remaining genes from T1L and T3D have superior oncolytic properties.

Three reassortants were chosen for testing of oncolytic properties relative to their parental viruses. Each of the reassortants , EB96, EB108 and EB146 posessed the M1 gene of T3D and were expected to preferentially replicate in cells that were damaged in their interferon response. These reassortants also possessed their L1, L3 and S2 genes of T1L that would be predicted to provide optimal replication abilities.

Oncolytic testing was performed by intranasal infection of 10^7 pfu of each virus into mice that possessed lung tumors derived form the CT26 colon tumor cell line fo Balb-C origin. Adult female BALB-C mice, 4-6 weeks old, were injected in the tail vein with 3×10^5 CT 26 on day 0 of the experiment. On day 7 groups of 3 mice were anaesthetized and infected with 10^7 pfu of virus in a 0.050 volume of culture medium. Mice were housed for an additional 6 days before euthanization with 90% CO₂/10% O₂. Lungs were removed, weighed, fixed in formalin and

photographed. One set of lungs was examined histopathologically by hematoxylin and eosin staining after paraffin embedding and sectioning.

The gross appearance of lungs after treatment showed that the untreated control lungs were heavily tumor laden having a pebbled surface appearance due to contiguous tumor nodules (Fig 3). These animals were in the terminal stages of cancer since one animal died at this time and the others were in respiratory distress. These lungs were 3 times heavier than uninfected balb-c lungs indicating the increased tumor mass approximated twice the mass of the lung tissue (Fig 4). Histologically these lungs were covered with a contiguous layer of tumor nodules and internal tumor masses seen as eosinophilic growths of cells (Fig 4 and 5). Infection with T1L virus resulted in a partial freeing of surface tumor growth observable on gross inspection that was also associated with a decrease in interior and surface nodules and a 20 % reduction in lung weight relative to untreated control (Fig 3, 4 and 5). T3D treatment was not as effective as T1L resulting in lungs that were only distinguishable from untreated controls by a slight (8 %) decrease in size but were similar in gross and microscopic appearance of tumors (Fig 3, 4 and 5). In dramatic contrast the EB96 reassortant virus cleared the lung of gross tumor mass on treatment (Fig 3). The lungs were of approximately normal weight having been freed of tumor masses (Fig 4). A small number of residual tumor cells remained at this time as detected by histological examination (Fig 5). The lungs were of normal size and appearance except for some circular patterns and dents on the lungs surface that presumably marked the location of prior tumor nodules. EB146 virus was not more effective at tumor lysis than the T3D parental virus (Fig 3, 4 and 5). Reassortant EB108 was partially effective at oncolysis producing results that were marginally better but similar than the T1L parental strain. On comparison of the genotypes of the reassortants it can be seen that the 3 reassortants possess 7 genome segments in common and thus differ in their L2, S3 and S4 genome segments indicating that the latter group of genes include important modulators of oncolysis. The EB96 reassortant is more effective than EB108 solely due to the nature of the S4 gene since these viruses only differ in the parental origin of this gene. This indicates that the T1L S4 gene conferred enhanced oncolytic properties relative to the T3D S4 gene. Since the S4 gene encodes the dsRNA binding protein that blocks PKR activation it is possible that the T1L S4 gene differs in this ability and thus, in concert with other combinations of T1L and T3D genome segments, controls oncolytic potential. In conclusion, the

dramatic increase in effectiveness of the EB96 reassortant at oncolysis, relative to the parental T1L and T3D viruses demonstrates the proof of principle that reassortants of reovirus with specific genotypes have enhanced and effective tumor lysis abilities in metastatic tumors in hosts with active immune responses. Table 6.

Table 6: Ranking of the ability of reovirus reassortants to lyse ct26 lung tumors. The relative weight of ct26 tumor bearing lungs relative to untreated control tumor bearing lungs are shown. The parental origin of genome segments are indicated as L for T1L and D for T3D.

VIRUS	TUMOR %	L1	L2	L3	M1	M2	M3	S1	S2	S3	S4	RANK
eb96	41	L	D	L	D	L	L	L	L	D	L	1
eb108	75	L	D	L	D	L	L	L	L	D	D	2
T1L	80	L	L	L	L	L	L	L	L	L	L	3
eb146	89	L	L	L	D	L	L	L	L	L	D	4
T3D	92	D	D	D	D	D	D	D	D	D	D	5

Experiment 6: Ability of T1L x T3D Reassortants to lyse tumors in vitro

A panel of tumor cell lines obtained from the NCI tumor panel (SF539, cns; SKMEL28, melanoma; HT29; NCI H23, nsc-lung; SW620, colon; DU145, prostate) were infected with the T1L, T3D, or the reassortants , EB96, EB108 and EB146 at an moi of 10 and were observed for cytopathic effect over a 5 day period. The ability to lyse tumor cells was scored visually on a scale of – to +++, where – indicates no difference from mock infected cells and +, ++, and +++ indicate 33 % cell destruction, 66 % cell destruction and complete lysis respectively. Although different tumor cell types differed in their susceptibility to lysis by different reovirus parents or reassortants the reassortants viruses were all as effective or more effective than the T3D parental virus at tumor cell lysis in vitro (Table 7).

Table 7: Cytopathology of reovirus T1L and T3D and reassortants
in different tumor cell lines

	Tumor cell line					
virus	SFS39 Cns	SKMEL28 melanoma	HT29	NCI H23 nsc-lung	SW620 colon	DU145 prostate
T1L	++	+++	++	+++	-	++
T3D	-	+++	+	++	-	+
EB96	++	+++	++	++	+	+
EB108	++	+++	++	++	+	+
EB146	++	+++	++	+++	+	++
RAS						

CLAIMS

What is claimed is:

1. A method of identifying a PKR sensitive virus comprising:
 - a) dividing a sample of a virus to be tested into a first portion and second portion;
 - b) contacting PKR +/+ cells with the first portion and contacting PKR -/- cells with the second portion, under conditions permitting growth of the virus in PKR -/- cells;
 - c) determining the rate of growth of the virus in the PKR +/+ cells and in the PKR -/- cells; and
 - d) comparing the growth rates from step c), wherein a higher rate of growth in the PKR -/- cells than in the PKR +/+ cells identifies the virus as PKR sensitive.
2. The method of claim 1, wherein the PKR cells are mouse embryo fibroblasts.
3. The method of claim 1, wherein the rate of growth of the virus is determined by a technique selected from the group consisting of plaque titer assay, antibody assay, and Western blot.
4. The method of claim 1, wherein the growth rate in PKR -/- cells is at least ten times higher than the growth rate in PKR +/+ cells.
5. A method of reducing the viability of a tumor cell, comprising administering to the tumor cell a virus identified as PKR-sensitive in the method of any one of claims 1-4.

6. A method of infecting a neoplasm in a mammal with a virus, comprising administering to the mammal a virus identified as PKR-sensitive in the method of any one of claims 1-4.
7. A method of treating a neoplasm in a mammal comprising administering to the mammal a therapeutically effective amount of a virus identified as PKR-sensitive in the method of any one of claims 1-4.
8. Use of the virus identified as PKR-sensitive in the method of any one of claims 1-4 in the manufacture of a medicament for reducing the viability of a tumor cell, infecting a neoplasm in a mammal or treating a neoplasm in a mammal.
9. The method or use of any one of claims 5-8, wherein the virus is administered by a route selected from the group consisting of intranasally, intratracheally, intravenously, intraperitoneally or intratumorally.
10. The method or use of any one of claims 5-9 wherein the virus is administered to a human or non-human mammal.
11. The method or use of claim 9 or 10 wherein the virus is administered at a dose of from 3×10^7 to 3×10^9 PFU/kg.

YIELD IN PKR-/- VS PKR+/- MEF

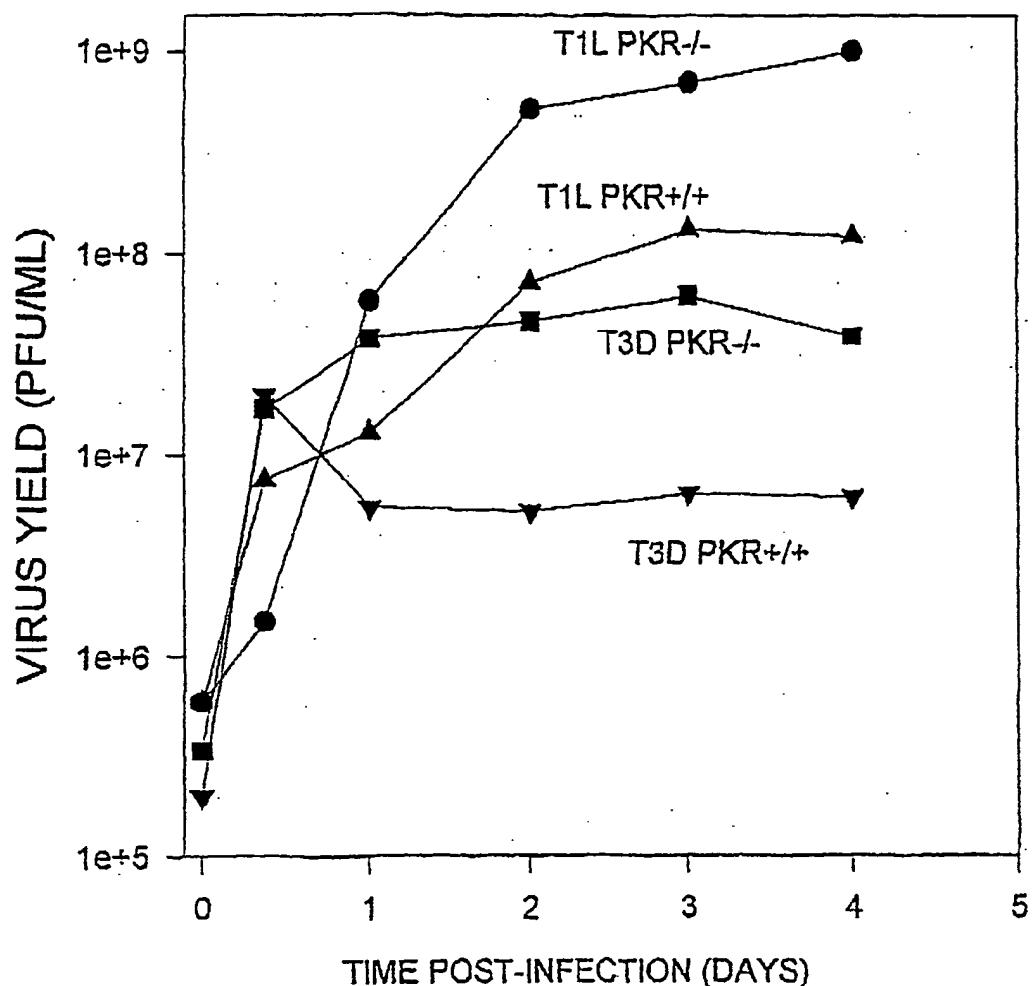


FIGURE 1

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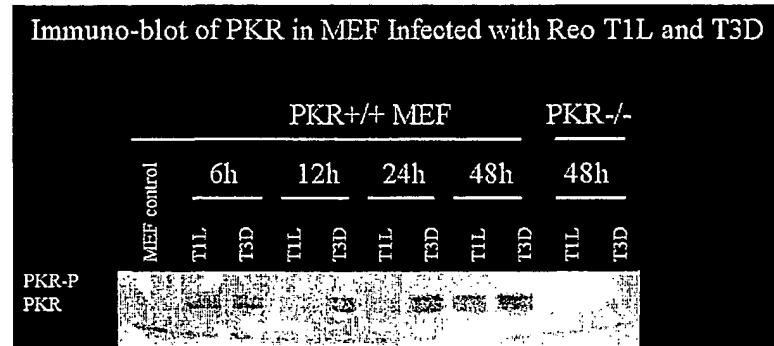


FIGURE 2

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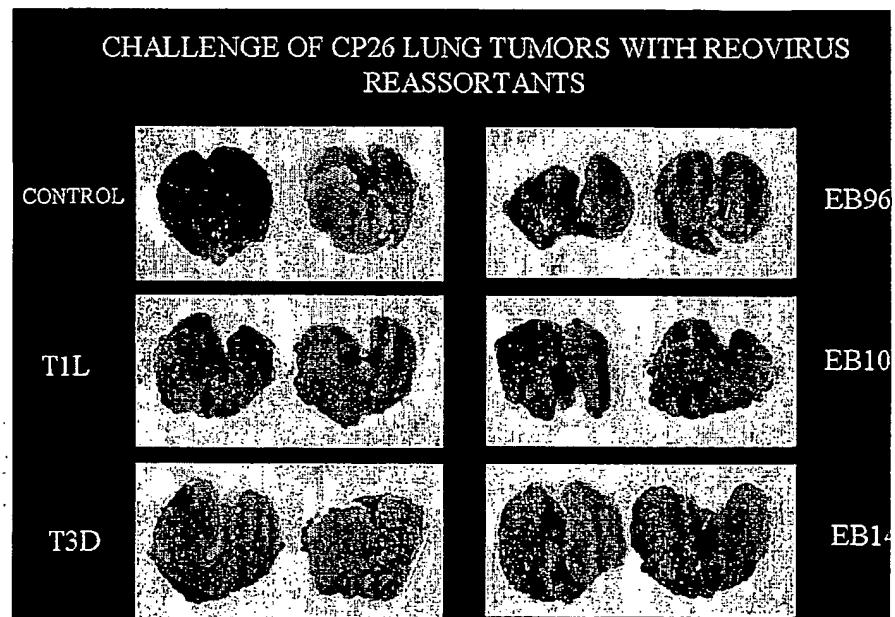


FIGURE 3

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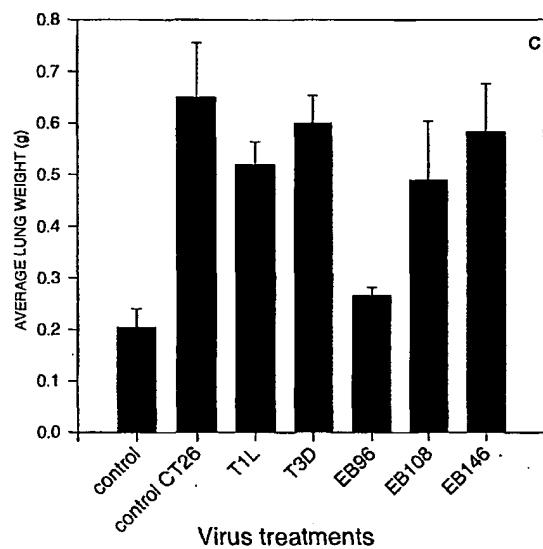
LUNG WEIGHT 6 DAYS POST TREATMENT

FIGURE 4
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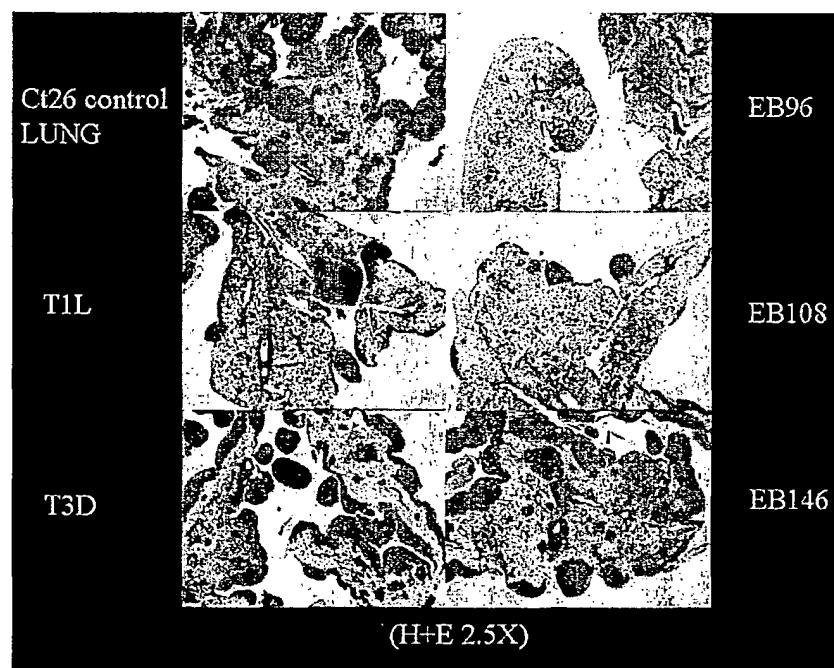


FIGURE 5

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